

FIG. 1A

FIG. 1B

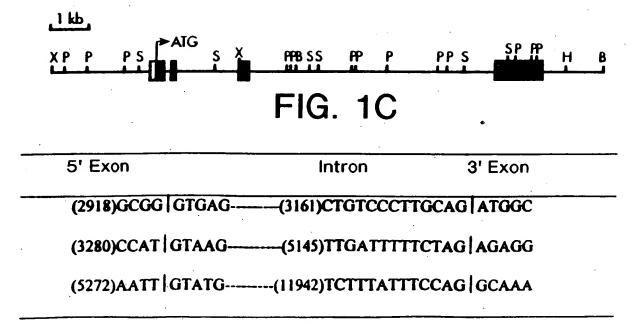


FIG. 1D



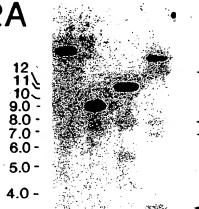
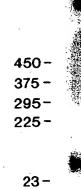


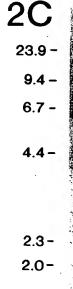
FIG. 2B



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9.5 -7.5 -

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9.5 -

7.5-

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1.35 -

FIG. 3A

FIG. 3B

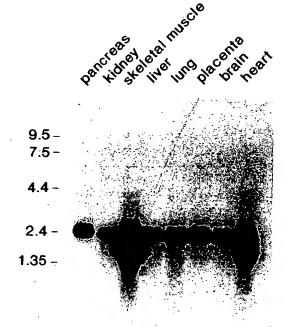
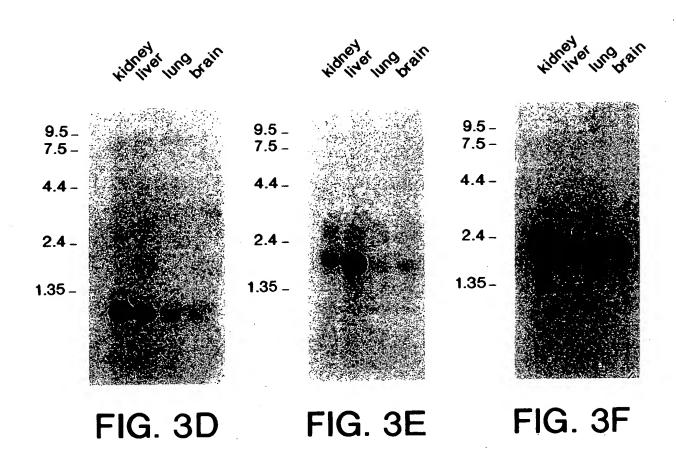


FIG. 3C

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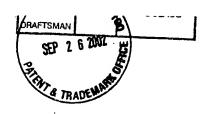
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# FIG. 4B-2

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tγ		PRO	. PRO	. PRO	. PRO	. PRO	. PRO	. PRO	. PRO	. PRO	. PRO	. PRO
Majority		1,49433	P41436	P41437	19251	U32373	U32974	U36842	45878	U45879	U45880	U45881
	0	H H	л П	<u>р</u> .	$\Box$					$\Box$	D	(EQ)
RDWEPGDDPWEEHAKWFPRCEFLLLAKGQEYVS	400	DEFY/Y	XDX 	RDEW-	SAEVI	PSIVIS	DEYIN	<b>DEY</b> IN	DEFIR	O EFFOR	DEVIN	PAYVIS
LAKG		1 2 2 2 2 2 2 2 2 2 2 2 2 3 3 3 3 3 3 3	了 次 6	17000000000000000000000000000000000000	SMX SI	LAKG	EOXG(	<b>DEXG</b>	RIXG(	RMXG(	EOKG(	LAKG
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XDW T	37		设	0	K K	S	Į.	Ď Ž	S	S S S S	Ŭ S	SI SI SI SI SI SI SI SI SI SI SI SI SI S
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GIG		DRN AFO	GYG	9000 0000	公気	X10	の国の		GNS	SKN SKN	6 B C	A SIGNATURE
Z		\$\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	线	刻	经	8	AL	AL	N X	XX	AL	ŎĶ Į



## FIG. 4A

10 20 30 40

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## FIG. 4B-1

-TPQELAKAGFYYLGRGDQVQCFACGGKLA Majority EEARLVTFQNWPD-AFI

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		. PRO	. PRO	. PRO	. PRO	. PRO	. PRO	. PRO	. PRO	. PRO	. PRO	. PRO	. PRO	IN. PRO
		L49433	L49441	P41436	P41437	U19251	U32373	~	U36842	U45878	U45879	U45880	U45881	SURVIV
•	250	SQKUIS	NGVIA	KVEIM	KVEIT	57,555	MAVIA	公公公尺尺	公安方面	公公内1S	公公内3S	2000万天	NGVIA	
	240	PICKNA (CEIA)	RLALAKOVA	RSDEVIROAFO	RGDEVINGA FC	KODÍTVOCESO	RLDHAKOVWO	I GOO O O O O O O O O O O O O O O O O O	A DO DO DO DE LO COMO DE LA COMO DEL COMO DE LA COMO DEL COMO DE LA COMO DEL COMO DEL COMO DEL COMO DEL COMO DEL COMO DE LA COMO DEL	PGTRAGAGAG	PODRIVACION	コロからなら知らり	RICHARCOWC	
	230	JARKSEY YEB	JAKKGEYYLLN	MAKINGEYYLG	MATASSELYYLG	15EAGEVE16	JAKÁGÉVYLIN	JAISAGLIVYITG	JAISTOCKTY VEG	JARAGEYYIG	JARAGEVYIE	ASKALVYKA	AKAGEYYLN	4A <u>ÉÁĞÉ</u> II
	220	1SPAE	IIII	LSPET	LEPSR	G18FCVI	ITPOA	LTPRE	IIPRD	LSPTDI	<u> </u>	17TPRE	IITPQA[	G-CAGTEER
	210	<b>  英色名的FLJTIY SIMが月-LSF1</b>	DANKINTEKDOON-PN	EDVELINTEER MOVE - SF1		<b>FEARTASERNMPEXYO</b>	$\overline{}$	<b>D</b>	D. D		EPEAREL LIXHIMAGE - LIFE	内内女女/大区S/DOS/WADDAIA		KDHKITSLEKKKKKH-LLK
		177	$\vdash$		$\infty$	ഗ	113	SO '	S O	SO.	∞.	യ	1 1 1	12

#### MWx10-3

200 -

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69 -

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Survivin

FIG. 4C-1

#### MWx10-3

200 -

97.4 -

69 -

46 -

30-

21.5-

14.3-

RblgG

FIG. 4C-2

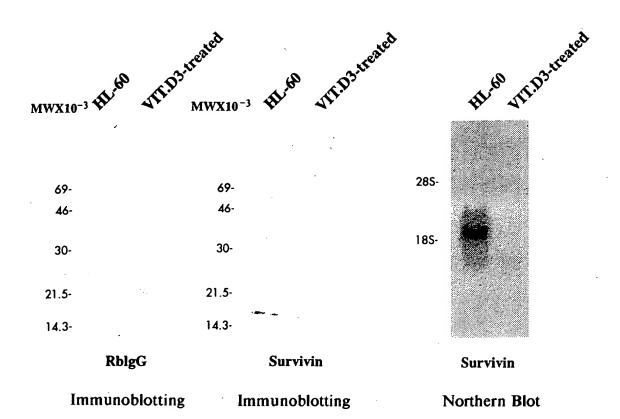


FIG. 5A FIG. 5B FIG. 5C

SCI 4.9 MIC



FIG. 6A



FIG. 6C

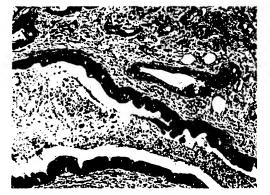


FIG. 6E

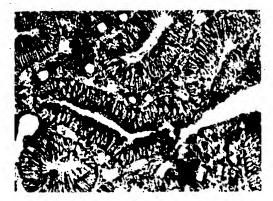


FIG. 6G

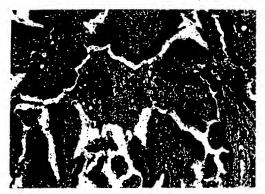


FIG. 6B

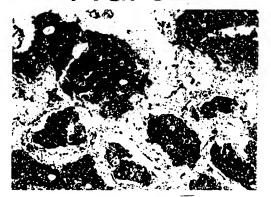


FIG. 6D

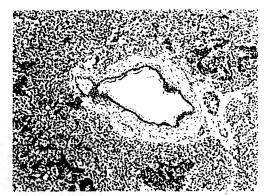


FIG. 6F



FIG. 6H

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MW x10<sup>-3</sup>

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69 -

46 -

30 -

21.5 -

14.3 -

FIG. 7A

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FIG. 7B-1

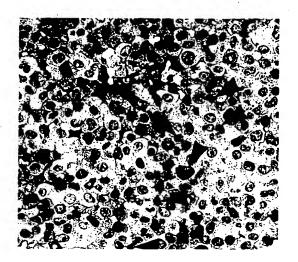


FIG. 7B-2

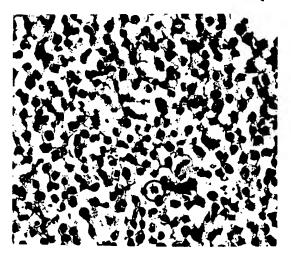


FIG. 7B-3

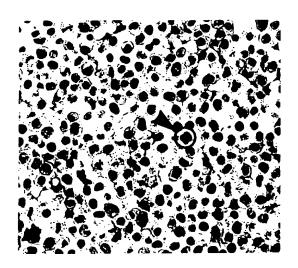


FIG. 7B-4

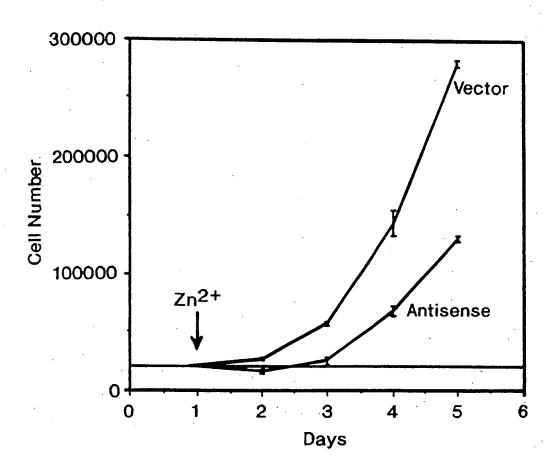


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FIG. 7C



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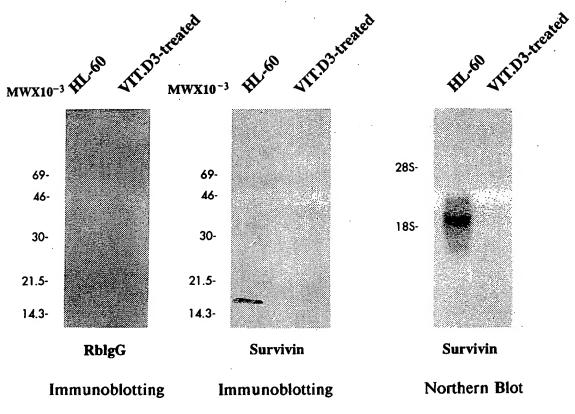


FIG. 8A FIG. 8B FIG. 8C

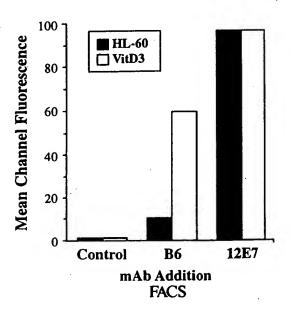
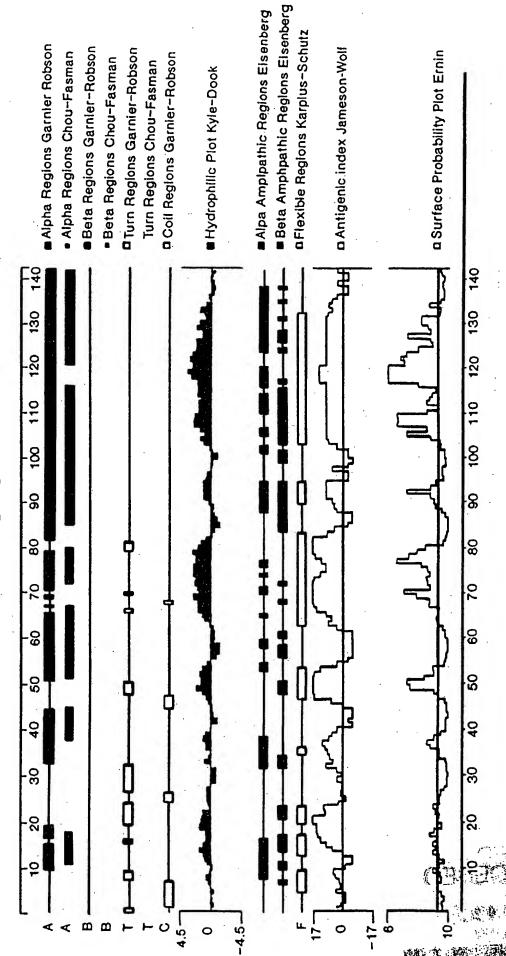


FIG. 8D







#### FIG. 10A

translation=MGAPTLPPAWQPFLKDHRISTFKNWPFLEGCACTPERMAEAGFIHCP TENEPDLAQCFFCFKELEGWEPDDDPIEEHKKHSSGCAFLSVKKQFEELTLGEFL KLDRERAKNKIAKETNNKKKEFEETAKKVRRAIEQLAAMD

•			•		
1	TCTAGACATG	CGGATATATT	CAAGCTGGGC	ACAGCACAGC	AGCCCCACCC
51	CAGGCAGCTT	GAAATCAGAG	CTGGGGTCCA	AAGGGACCAC	ACCCCGAGGG
101	ACTGTGTGGG	GGTCGGGGCA	CACAGGCCAC	TGCTTCCCCC	CGTCTTTCTC
151	AGCCATTCCT	GAAGTCAGCC	TCACTCTGCT	TCTCAGGGAT	TTCAAATGTG
201	CAGAGACTCT	GGCACTTTTG	TAGAAGCCCC	TTCTGGTCCT	AACTTACACC
251	TGGATGCTGT	GGGGCTGCAG	CTGCTGCTCG	GGCTCGGGAG	GATGCTGGGG
301	GCCCGGTGCC	CATGAGCTTT	TGAAGCTCCT	GGAACTCGGT	TTTGAGGGTG
351	TTCAGGTCCA	GGTGGACACC	TGGGCTGTCC	TTGTCCATGC	ATTTGATGAC
401	ATTGTGTGCA	GAAGTGAAAA	GGAGTTAGGC	CGGGCATGCT	GGCTTATGCC
451	TGTAATCCCA	GCACTTTGGG	AGGCTGAGGC	GGGTGGATCA	CGAGGTCAGG
501	AGTTCAATAC	CAGCCTGGCC	AAGATGGTGA	AACCCCGTCT	CTACTAAAAA
551	ТАСАААААА	TTAGCCGGGC	ATGGTGGCGG	GCGCATGTAA	TCCCAGCTAC
601	TGGGGGGCT	GAGGCAGAGA	ATTGCTGGAA	CCCAGGAGAT	GGAGGTTGCA
651	GTGAGCCAAG	ATTGTGCCAC	TGCACTGCAC	TCCAGCCTGG	CGACAGAGCA
701	AGACTCTGTC	TCAAAAAAAA	AAAAAAAAAG	TGAAAAGGAG	TTGTTCCTTT
751	CCTCCCTCCT	GAGGGCAGGC	AACTGCTGCG	GTTGCCAGTG	GAGGTGGTGC
801	GTCCTTGGTC	TGTGCCTGGG	GGCCACCCCA	GCAGAGGCCA	TGGTGGTGCC
851	AGGGCCCGGT	TAGCGAGCCA	ATCAGCAGGA	CCCAGGGGCG	ACCTGCCAAA
901	GTCAACTGGA	TTTGATAACT	GCAGCGAAGT	TAAGTTTCCT	GATTTTGATG
951	ATTGTGTTGT	GGTTGTGTAA	GAGAATGAAG	TATTTCGGGG	TAGTATGGTA
1001	ATGCCTTCAA	CTTACAAACG	GTTCAGGTAA	ACCACCCATA	TACATACATA
1051	TACATGCATG	TGATATATAC	ACATACAGGG	ATGTGTGTGT	GTTCACATAT
1101	ATGAGGGGAG	AGAGACTAGG	GGAGAGAAAG	TAGGTTGGGG	AGAGGGAGAG
1151	AGAAAGGAAA	ACAGGAGACA	GAGAGAGAGC	GGGGAGTAGA	GAGAGGGAAG
1201	GGGTAAGAGA	GGGAGAGGAG	GAGAGAAAGG	GAGGAAGAAG	CAGAGAGTGA
1251	ATGTTAAAGG	AAACAGGCAA	AACATAAACA	GAAAATCTGG	GTGAAGGGTA
1301	TATGAGTATT	CTTTGTACTA	TTCTTGCAAT	TATCTTTTAT	TTAAATTGAC
1351	ATCGGGCCGG	GCGCAGTGGC	TCACATCTGT	AATCCCAGCA	CTTTGGGAGG
1401		•			AGCCTGGCAA
1451	ACATGGTGAA	ACCCCATCTC	TACTAAAAAT	ACAAAAATTA	GCCTGGTGTG
1501	GTGGTGCATG	CCTTTAATCT	CAGCTACTCG	GGAGGCTGAG	GCAGGAGAAT
1551					GATCATGCCA
1601	•				AATAAATAAA
1651					GCGGGGTGGG
1701	•			•	TAAACCCCAA
1751	· ·				GGAGGGCTAA
1801	•			AACCAGAGAA	
1851					CCAAAGCAGA
1901					TGTTGGTGGG
1951					GCACAGGCCC
2001					ATCACGGTAG
2051					CTGGAAACCC
2101					TCTGCTGCAC
2151		•	•		GAGTAGATGC.

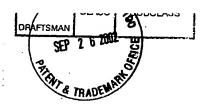


### FIG. 10B

2201	TTTTTGCAGA	GGTGGCACCC	TGTAAAGCTC	TCCTGTCTGA	CTTTTTTTT
2251	TTTTTTAGAC	TGAGTTTTGC	TCTTGTTGCC	TAGGCTGGAG	TGCAATGGCA
2301	CAATCTCAGC	TCACTGCACC	CTCTGCCTCC	CGGGTTCAAG	CGATTCTCCT
2351	GCCTCAGCCT	CCCGAGTAGT	TGGGATTACA	GGCATGCACC	ACCACGCCCA
2401	GCTAATTTTT	GTATTTTTAG	TAGAGACAAG	GTTTCACCGT	GATGGCCAGG
2451	CTGGTCTTGA	ACTCCAGGAC	TCAAGTGATG	CTCCTGCCTA	GGCCTCTCAA
2501	AGTGTTGGGA	TTACAGGCGT	GAGCCACTGC	ACCCGGCCTG	CACGCGTTCT
2551	TTGAAAGCAG	TCGAGGGGGC	GCTAGGTGTG	GGCAGGGACG	AGCTGGCGCG
2601	GCGTCGCTGG	GTGCACCGCG	ACCACGGGCA	GAGCCACGCG	GCGGGAGGAC
2651	TACAACTCCC	GGCACACCCC	GCGCCGCCCC	GCCTCTACTC	CCAGAAGGCC
2701	GCGGGGGGTG	GACCGCCTAA	GAGGGCGTGC	GCTCCCGACA	TGCCCCGCGG
2751	CGCGCCATTA	ACCGCCAGAT	TTGAATCGCG	GGACCCGTTG	GCAGAGGTGG
		>Sta			
2801	CGGCGGCGGC	ATGGGTGCCC	CGACGTTGCC	CCCTGCCTGG	CAGCCCTTTC
2851	TCAAGGACCA	CCGCATCTCT	ACATTCAAGA	ACTGGCCCTT	CTTGGAGGGC
2901	TGCGCCTGCA	CCCCGGAGCG	GGTGAGACTG	CCCGGCCTCC	TGGGGTCCCC
2951	CACGCCCGCC	TTGCCCTGTC	CCTAGCGAGG	CCACTGTGAC	TGGGCCTCGG
3001				GTCCCCAGCG	
3051	GGCTGGGCCC	CTTGGGTCCA	GGCCGGCCTC	CCCTCCCTGC	TTTGTCCCCA
3101				CCGGGCTGCC	•
3151				AGGCTGGCTT	
3201				TGTTTCTTCT	
3251	GCTGGAAGGC	TGGGAGCCAG	ATGACGACCC	CATGTAAGTC	TTCTCTGGCC
3301				TGTCAAAAGA	•
3351				CACCCTCATT	·
3401				GCTGGTGCCT	
3451				ATGCCTTGGG	
3501				TACATCAGCA	
3551	•			TTTTTGAGAT	
3601	ACAGTTTAAT	TGAAATATAA	CCTACACAGC	ACAAAAATTA	CCCTTTGAAA
3651	GTGTGCACTT	CACACTTTCG	GAGGCTGAGG	CGGGCGGATC	ACCTGAGGTC
3701	AGGAGTTCAA	GACCTGCCTG	GCCAACTTGG	CGAAACCCCG	TCTCTACTAA
3751	AAATACAAAA	ATTAGCCGGG	CATGGTAGCG	CACGCCCGTA	ATCCCAGCTA
3801	CTCGGGAGGC	TAAGGCAGGA	GAATCGCTTG	AACCTGGGAG	GCGGAGGTTG
3851	CAGTGAGCCG	AGATTGTGCC	AATGCACTCC	AGCCTCGGCG	ACAGAGCGAG
3901	•		•	AAAAAAAGAA	
3951			•	CAAGATGCCT	
4001	AATGAAATTC	TGTACTCGGA	TGGTATCTGT	CTTTCCACAC	TGTAATGCCA
4051			•	TTCAGTTGCT	•
4101				TTTGCTATGG	
4151				TGGTATTCTT	
4201		·		TTACAACTGG	
4251				TGCAGTTCTG	
4301			•		AGTGTGAGCT
4351				ATGCCCTTCT	
4401				CGCCTGTAAT	
4451				CAGGAGATCG	
4501				AAAATACAAA	

## FIG. 10C

4551	GGCGTGGTGG	TGGGCGCCTG	TAGTCCCAGC	TACTCGGGAG	GCTGAGGCAG
4601	GAGAATGGCG	TGAACCCAGG	AGGCGGAGCT	TGCAGTGAGC	CGAGATTGCA
4651	CCACTGCACT	CCAGCCTGGG	CGACAGAATG	AGACTCCGTC	TCAAAAAAAA
4701	AAAAAAAAGA	AAAAAATCTT	TACAGTGGAT	TACATAACAA	TTCCAGTGAA
4751	ATGAAATTAC	TTCAAACAGT	TCCTTGAGAA	TGTTGGAGGG	ATTTGACATG
4801	TAATTCCTTT	GGACATATAC	CATGTAACAC	TTTTCCAACT	AATTGCTAAG
4851	GAAGTCCAGA	TAAAATAGAT	ACATTAGCCA	CACAGATGTG	GGGGGAGATG
4901	TCCACAGGGA	GAGAGAAGGT	GCTAAGAGGT	GCCATATGGG	AATGTGGCTT
4951	GGGCAAAGCA	CTGATGCCAT	CAACTTCAGA	CTTGACGTCT	TACTCCTGAG
5001	GCAGAGCAGG	GTGTGCCTGT	GGAGGGCGTG	GGGAGGTGGC	CCGTGGGGAG
5051	TGGACTGCCG	CTTTAATCCC	TTCAGCTGCC	TTTCCGCTGT	TGTTTTGATT
5101	TTTCTAGAGA	GGAACATAAA	AAGCATTCGT	CCGGTTGCGC	TTTCCTTTCT
5151	GTCAAGAAGC	AGTTTGAAGA	ATTAACCCTT	GGTGAATTTT	TGAAACTGGA
5201	CAGAGAAAGA	· <u> </u>			AAGAACTGCT
5251	CAAACCCTGT	TCAATGTCTT	TAGCACTAAA	CTACCTAGTC	CCTCAAAGGG
5301	ACTCTGTGTT	TTCCTCAGGA		TTTTTTTCT	GAGATAGAGT
5351	TTCACTCTTG	TTGCCCAGGC		TGGTGCAATC	TTGGCTCACT
5401	GCAACCTCTG	CCTCTCGGGT	TCAAGTGATT	CTCCTGCCTC	
5451	GTAACTGGGA	TTACAGGGAA	GTGCCACCAC	ACCCAGCTAA	TTTTTGTATT
5501				CCAGGCTGGT	
5551		ATTCGCCCAC	CTTGGCCTCC	CAAAGTGCTG	
5601	CGTGAACCAC		TTTTTTTTT	TTGTTCTGAG	ACACAGTTTC
5651	ACTCTGTTAC		GTAGGGTGGC	CTGATCTCGG	ATCACTGCAA
5701	CCTCCGCCTC			TGCTTCAGCC	•
5751		•	CACCACACCC		TGTATTTTTG
5801	•		•	GCTGGTTTTG	AACTCCTGAC
5851	CTCAGGTGAT			AAGTGCTGAG	
5901	TGAGCCACCA	CACCTGGCCT	CAGGAAGTAT	TTTTATTTT	AAATTTATTT
5951	ATTTATTTGA	GATGGAGTCT	TGCTCTGTCG	CCCAGGCTAG	AGTGCAGCGA
6001	CGGGATCTCG	GCTCACTGCA	AGCTCCGCCC	CCCAGGTTCA	AGCCATTCTC
6051	CTGCCTCAGC	CTCCCGAGTA	GCTGGGACTA	CAGGCGCCCG	CCACCACACC
6101	CGGCTAATTT	TTTTGTATTT	TTAGTAGAGA	CGGGTTTTCA	CCGTGTTAGC
6151	CAGGAGGGTC	TTGATCTCCT	GACCTCGTGA	TCTGCCTGCC	TCGGCCTCCC
6201				ACACCCGGCT	
6251				GGCTGCAGTG	
6301	CCATAGCTCA	CTGCAGCCTC	GAACTCCTGA	GCTCAAGTGA	TCCTCCCACC
6351				GCACCCCACC	
6401				CATAGAGATG	
6451	GTGTTGTCCA	GGCTGGTCTT	GAACTCCTGA	GCTCACGGGA	TCCTTTTGCC
6501				ATGAGCCACC	
6551	AGGAATCATT				
6601				ATAAGTAAAT	
6651					ATTTTTTT
6701				TGGAGTGCAG	
6751				TCAAGCGATT	·
6801	AGCCTCCCGA				
6851					AAGCTGGTCT
6901	•				GAGTGCTGGG



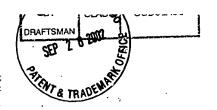
### FIG. 10D

6951	ATTACAGGCA	TGAGCCACCA	TGCGTGGTCT	TTTTAAAATT	TTTTGATTTT
7001	TTTTTTTTT	GAGACAGAGC	CTTGCTCTGT	CGCCCAGGCI	GGAGTGCAGT
7051	GGCACGATCT	CAGCTCACTA	CAAGCTCCGC	CTCCCGGGTT	CACGCCATTC
7101	TTCTGCCTCA	GCCTCCTGAG	TAGCTGGGAC	TACAGGTGCC	CACCACCACG
7151	CCTGGCTAAT	TTTTTTTGGT	ATTTTTATTA	GAGACAAGGT	TTCATCATGT
7201	TGGCCAGGCT		TCCTGACCTC	AAGTGATCTG	CCTGCCTCGG
7251	CCTCCCAAAG	CGCTGAGATT	ACAGGTGTGA	TCTACTGCGC	CAGGCCTGGG
7301	CGTCATATAT	TCTTATTTGC	TAAGTCTGGC	AGCCCCACAC	AGAATAAGTA
7351	CTGGGGGATT		GTAGCAAAGC	CCTGGGTGGA	GAGTCAGGAG
7401	ATGTTGTAGT	TCTGTCTCTG	CCACTTGCAG	ACTTTGAGTT	TAAGCCAGTC
7451	GTGCTCATGC	TTTCCTTGCT	AAATAGAGGT	TAGACCCCCT	ATCCCATGGT
7501	TTCTCAGGTT	GCTTTTCAGC	TTGAAAATTG	TATTCCTTTG	TAGAGATCAG
7551	CGTAAAATAA		ATATGTGGCT	TTATTTTAAT	TTGAGACAGA
7601	GTGTCACTCA	GTCGCCCAGG	CTGGAGTGTG	GTGGTGCGAT	CTTGGCTCAC
7651	TGCGACCTCC	ACCTCCCAGG	TTCAAGCGAT	TCTCGTGCCT	CAGGCTCCCA
7701	AGTAGCTGAG	ATTATAGGTG	TGTGCCACCA	GGCCCAGCTA	ACTTTTGTAT
7751	TTTTAGTAGA	GACAGGGTTT	TGCCATGTTG	GCTAAGCTGG	TCTCGAACTC
7801	CTGGCCTCAA	GTGATCTGCC	CGCCTTGGCA	TCCCAAAGTG	CTGGGATTAC
7851	AGGTGTGAAC	CACCACACCT	GGCCTCAATA	TAGTGGCTTT	TAAGTGCTAA
7901	GGACTGAGAT	TGTGTTTTGT	CAGGAAGAGG	CCAGTTGTGG	GTGAAGCATG
7951	CTGTGAGAGA	GCTTGTCACC	TGGTTGAGGT	TGTGGGAGCT	GCAGCGTGGG
8001	AACTGGAAAG	TGGGCTGGGG	ATCATCTTTT	TCCAGGTCAG	GGGTCAGCCA
8051	GCTTTTCTGC	AGCGTGCCAT	AGACCATCTC	TTAGCCCTCG	TGGGTCAGAG
8101	TCTCTGTTGC	ATATTGTCTT	TTGTTGTTTT	TCACAACCTT	TTAGAAACAT
8151	AAAAAGCATT	CTTAGCCCGT	GGGCTGGACA	AAAAAAGGCC	ATGACGGGCT
8201	GTATGGATTT	GGCCCAGCAG	GCCCTTGCTT	GCCAAGCCCT	GTTTTAGACA
8251	AGGAGCAGCT	TGTGTGCCTG	GAACCATCAT	GGGCACAGGG	GAGGAGCAGA
8301	GTGGATGTGG	AGGTGTGAGC	TGGAAACCAG	GTCCCAGAGC	GCTGAGAAAG
8351	ACAGAGGGTT		CAAGTAGAGC		TGACACCATC
8401	CAGTTCCAGA			ACGCTGCGGG	
8451	CTAGGGTTAC	•			TCCACTCACC
8501				TTTCAGGGCC	<del>-</del>
8551				GGCCGAGGCG	
8601				ACATGGTGAA	
8651				GGTGGCGGC	· ·
8701				TGGCGTGAAC	
8751				GCACTCCAGC	
8801				AAAGTAGGCT	
8851	GTGAGCTGAA				
8901	GACCCCTCGG	TCTCTATCTC	CTGATAGTCA	GACCCAGCCA	CACTGGAAAG
8951	AGGGGAGACA	TTACAGCCTG	CGAGAAAAGT	AGGGAGATTT	AAAAACTGCT
9001	TGGCTTTTAT	TTTGAACTGT	TTTTTTTGTT	TGTTTGTTTT	CCCCAATTCA
9051	GAATACAGAA				
9101					TACTCTGTCA
9151	CCCAGGCTGA				
9201	CTGGGCTCAA				
9251	AGGTGCGTGT				
9301	GGTGATGCTG	GGGATTCCCA	AGATCCCAGA	TTTGATGGCA	GGATGCCCCT.
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### FIG. 10E

9351	GTCTGCTGCC	TTGCCAGGGT	GCCAGGAGGG	CGCTGCTGTG	GAAGCTGAGG
9401	CCCGGCCATC	CAGGGCGATG	CATTGGGCGC	TGATTCTTGT	TCCTGCTGCT
9451	GCCTCGGTGC	TTAGCTTTTG	AAACAATGAA	ATAAATTAGA	ACCAGTGTGA
9501	AAATCGATCA	GGGAATAAAT	TTAATGTGGA	AATAAACTGA	ACAACTTAGT
9551	TCTTCATAAG	AGTTTACTTG	GTAAATACTT	GTGATGAGGA	CAAAACGAAG
9601	CACTAGAAGG	AGAGGCGAGT	TGTAGACCTG	GGTGGCAGGA	GTGTTTTGTT
9651	TGTTTTCTTT	GGCAGGGTCT	TGCTCTGTTG	CTCAGGCTGG	AGTACAGTGG
9701	CACAATCACA	GCTCACTATA	GCCTCGACCT	CCTGGACTCA	AGCAATCCTC
9751.	CTGCCTCAGC	CTCCCAGTAG	CTGGGACTAC	AGGCGCATGC	CACCATGCCT
9801	GGCTAATTTT	AAATTTTTTT	TTTTCTCTTT	TTTGAGATGG	AATCTCACTC
9851	TGTCGCCCAG	GCTGGAGTGC	AGTGGCGTGA	TCTCGGCTGA	CGGCAAGCTC
9901	CGCCTCCCAG	GTTCACTCCA	TTCGCCTGCC	TCAGCCTCCC	AAGTAGCTGG
9951	GACTACAGGC	GCTGGGATTA	CAAACCCAAA	CCCAAAGTGC	TGGGATTACA
10001	GGCGTGAGCC	ACTGCACCCG	GCCTGTTTTG	TCTTTCAATA	GCAAGAGTTG
10151	TGTTTGCTTC	GCCCCTACCT	TTAGTGGAAA	AATGTATAAA	ATGGAGATAT
10201	TGACCTCCAC	ATTGGGGTGG	TTAAATTATA	GCATGTATGC	AAAGGAGCTT
10251	CGCTAATTTA	AGGCTTTTTT	GAAAGAGAAG	AAACTGAATA	ATCCATGTGT
10301	GTATATATAT	TTTAAAAGCC	ATGGTCATCT	TTCCATATCA	GTAAAGCTGA
10351	GGCTCCCTGG	GACTGCAGAG	TTGTCCATCA	CAGTCCATTA	TAAGTGCGCT
10401	GCTGGGCCAG	GTGCAGTGGC	TTGTGCCTGA	ATCCCAGCAC	TTTGGGAGGC
10451	CAAGGCAGGA	GGATTCATTG	AGCCCAGGAG	TTTTGAGGCG	AGCCTGGGCA
10501	ATGTGGCCAG	ACCTCATCTC	TTCAAAAAAT	ACACAAAAAA	TTAGCCAGGC
10551	ATGGTGGCAC	GTGCCTGTAG	TCTCAGCTAC	TCAGGAGGCT	GAGGTGGGAG
10601	GATCACTTTG	AGCCTTGCAG	GTCAAAGCTG	CAGTAAGCCA	TGATCTTGCC
10651	ACTGCATTCC	AGCCTGGATG	ACAGAGCGAG	ACCCTGTCTC	AAAAAAAA
10701	AAAAACCAAA	CGGTGCACTG	TTTTCTTTTT	TCTTATCAAT	TTATTATTTT
10751	TAAATTAAAT	TTTCTTTTAA	TAATTTATAA	ATTATAAATT	TATATTAAAA
10801	AATGACAAAT	TTTTATTACT	TATACATGAG	GTAAAACTTA	GGATATATAA
10851	AGTACATATT	GAAAAGTAAT	TTTTTGGCTG	GCACAGTGGC	TCACACCTGT
10901	AATCCCAGCA	CTTTGGGAGG	CCGTGGCGGG	CAGATCACAT	GAGATCATGA
10951	GTTCGAGACC	AACCTGACCA	ACATGGAGAG	ACCCCATCTC	TACTAAAAAT
11001	ACAAAATTAG	CCGGGGTGGT	GGCGCATGCC	TGTAATCCCA	GCTACTCGGG
11051	AGGCTGAGGC	AGGAGAATCT	CTTGAACCCG	GGAGGCAGAG	GTTGCGGTGA
11101	GCCAAGATCG	TGCCTTTGCA	CACCAGCCTA	GGCAACAAGA	GCGAAAGTCC
11151	GTCTCAAAAA	AAAAGTAATT	TTTTTTAAGT	TAACCTCTGT	CAGCAAACAA
11201	ATTTAACCCA	ATAAAGGTCT	TTGTTTTTTA	ATGTAGTAGA	GGAGTTAGGG
11251	TTTATAAAAA	ATATGGTAGG	GAAGGGGGTC	CCTGGATTTG	CTAATGTGAT
11301	TGTCATTTGC	CCCTTAGGAG	AGAGCTCTGT	TAGCAGAATG	AAAAAATTGG
11351	AAGCCAGATT	CAGGGAGGGA	CTGGAAGCAA	AAGAATTTCT	GTTCGAGGAA
11401	GAGCCTGATG	TTTGCCAGGG	TCTGTTTAAC	TGGACATGAA	GAGGAAGGCT
11451				AGGTAGGGCA	
11501				CTAGACTAGC	
11551				TACTTTGAGT	
11601	•			TAGAGATATG	
11651				AATGGAGGAA	•
11701		•		AGTGTTTCCT	.,1
11751				TGAGCTGAAT	
11801				AGTGCCCAGT	
•	· - <del>*</del>				= <del></del>



## FIG. 10F

	11851	GCTCTCTCAG	TGTTCCCTGA	TTGTTTTTC	CTTTGTCATC	TTATCTACAG
	11901	GATGTGACTG	GGAAGCTCTG	GTTTCAGTGT	CATGTGTCTA	TTCTTTATTT
	11951	CCAGGCAAAG	GAAACCAACA	ATAAGAAGAA	AGAATTTGAG	GAAACTGCGA
	12001	AGAAAGTGCG	CCGTGCCATC	GAGCAGCTGG	CTGCCATGGA	TTGAGGCCTC
	12051	TGGCCGGAGC	TGCCTGGTCC	CAGAGTGGCT	GCACCACTTC	CAGGGTTTAT
	12101	TCCCTGGTGC	CACCAGCCTT	CCTGTGGGCC	CCTTAGCAAT	GTCTTAGGAA
	12151	AGGAGATCAA	CATTTTCAAA	TTAGATGTTT	CAACTGTGCT	CCTGTTTTGT
	12201	CTTGAAAGTG	GCACCAGAGG	TGCTTCTGCC	TGTGCAGCGG	GTGCTGCTGG
	12251	TAACAGTGGC	TGCTTCTCTC	TCTCTCTCTC	TTTTTTGGGG	GCTCATTTTT
	12301	GCTGTTTTGA	TTCCCGGGCT	TACCAGGTGA	GAAGTGAGGG	AGGAAGAAGG
	12351	CAGTGTCCCT	TTTGCTAGAG	CTGACAGCTT	TGTTCGCGTG	GGCAGAGCCT
	12401	TCCACAGTGA	ATGTGTCTGG	ACCTCATGTT	GTTGAGGCTG	TCACAGTCCT
	12451	GAGTGTGGAC	TTGGCAGGTG	CCTGTTGAAT	CTGAGCTGCA	GGTTCCTTAT
	12501	CTGTCACACC	TGTGCCTCCT	CAGAGGACAG	TTTTTTTTTTT	GTTGTGTTTT
	12551	TTTGTTTTTT	TTTTTTGGTA	GATGCATGAC	TTGTGTGTGA	TGAGAGAATG
	12601	GAGACAGAGT	CCCTGGCTCC		AACAACATGG	
	12651	TTGTTTGAAT			CAAACTACAA	
	12701				CGGGGTGAAC	
	12751				TCTGGCAGAT	,
	12801			•	AGCCGCGGGG	-
	12851				TAAATCCTTT	• • • • • • • • • • • • • • • • • • •
	12901	TGGCTCGATG			TGCTGCAGGC	CGTGTGTCTG
	12951				CCACACGGGG	
	13001	GTCCGCCCAG			CAGCAGCTCC	5
	13051	AAGTCTGGCG			GCCCTCCTCC	
	13101	GCTGCAGGGT			GAAACCTCTG	
	13151	CGGCTGTTCC			ATTTCAAACA	
	13201	CCCTACTGGG	TTTTTAAAAT		TTTCATCGTC	
	13251				GTGAGGATGA	
	13301				GAGCCACGAA	
	13351				AAACAACTGA	•
	13401				CGGCCTGAGA	
	13451				TTTGTTAGCA	
	13501			•	TTTTTGATAT	•
	13551			•	TTGCTTAGTA	
	13601				TCAACTCCTT	
	13651					GGCAGGGCTG
	13701	•		•	CACGGCCTTT	
	13751					AGCTTCCTCG
	13801				•	CCGCCTAGAC
	13851	·				CTCTAAGTTG
	13901					TATGGGTGAG
	•					
	13951 14001		•			CCTGGAAAGT CAAACTGGAG
	14001	,	•			CAAACTGGAG
	14101					CTTTTTTTT
•	14151					TGACCACAGC
	14201					ACTTTAGCCT
	T-4 % O T	* CUCCOCWAC	CICHMMIICI	TACATICAMA	TOWACCICCC	WCTTTWGCCT.

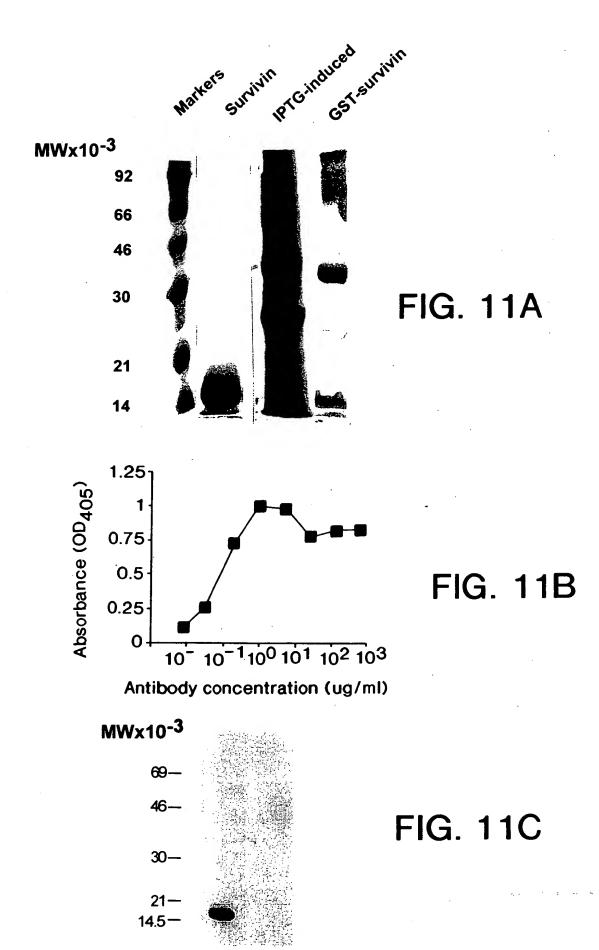


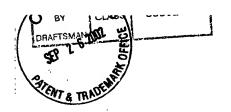
#### FIG. 10G

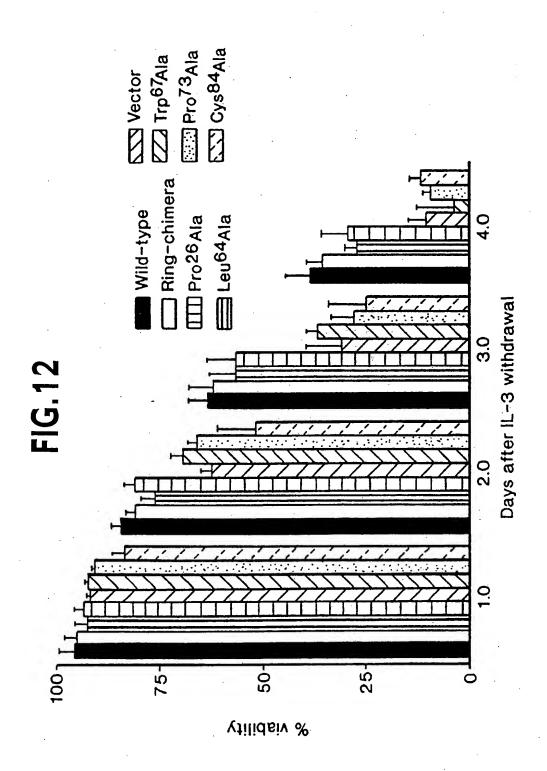
14251	CCCAAGTAGC	TGGGACTACA	GGCGCACGCC	ATCACACCCG	GCTAATTGAA
14301				ACTTTGTTGC	
14351				GCTTCAGCGT	
14401	GGTATTATAG	GCGTGAGCCA	CTGGGCCTGA	CCTAGCTACC	ATTTTTTAAT
14451				ACTTGTCCAG	
14501	TAAGTAACTT	TTAGAGCTGG	GATTTGAACC.	CAGGCAATCT	GGCTCCAGAG
14551	CTGGGCCCTC	ACTGCTGAAG	GACACTGTCA	GCTTGGGAGG	GTGGCTATGG
14601	TCGGCTGTCT	GATTCTAGGG	AGTGAGGGCT	GTCTTTAAAG	CACCCCATTC
14651				TCATATGGAG	
	CCTCCCCAAG				
14751	TGAAATGAAA	ATTCACAGGA	AGTTGTAAGG	CTAGTACAGG	GGATCC

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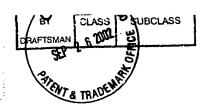






#### NEW TOTAL





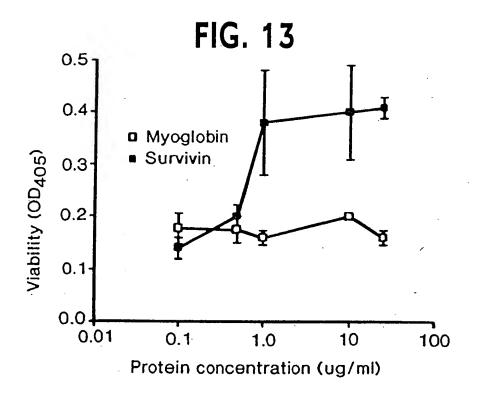
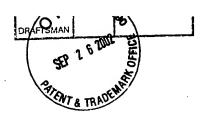
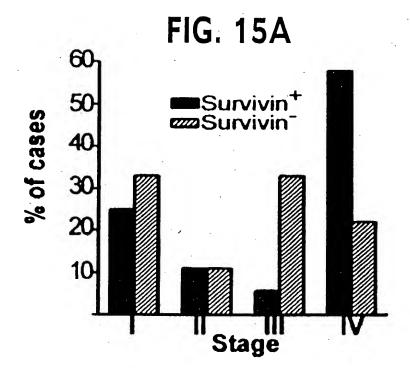


FIG. 14B **FIG. 14A** Prognostic factors Histology (Shimada class.) 757 75 n=72 % of survivin positive cases Survivin 50 % of cases 50 Survivin p=0.0025 25 Favorable Unfavorable 0 1 2 3 Number of negative prognostic factors (age, n-myc amplification Containing at least one widespread disease) negative prognostic factor





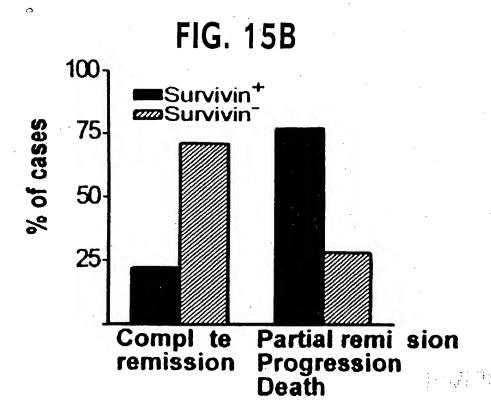
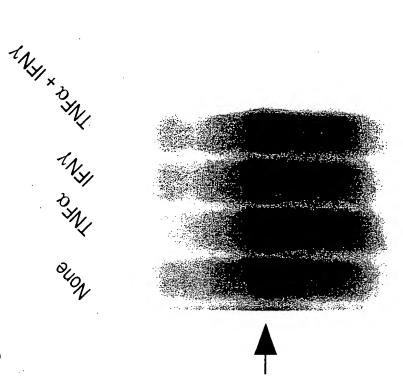


FIG. 16



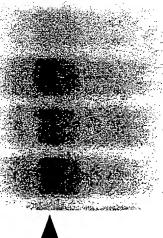




FIG. 17

